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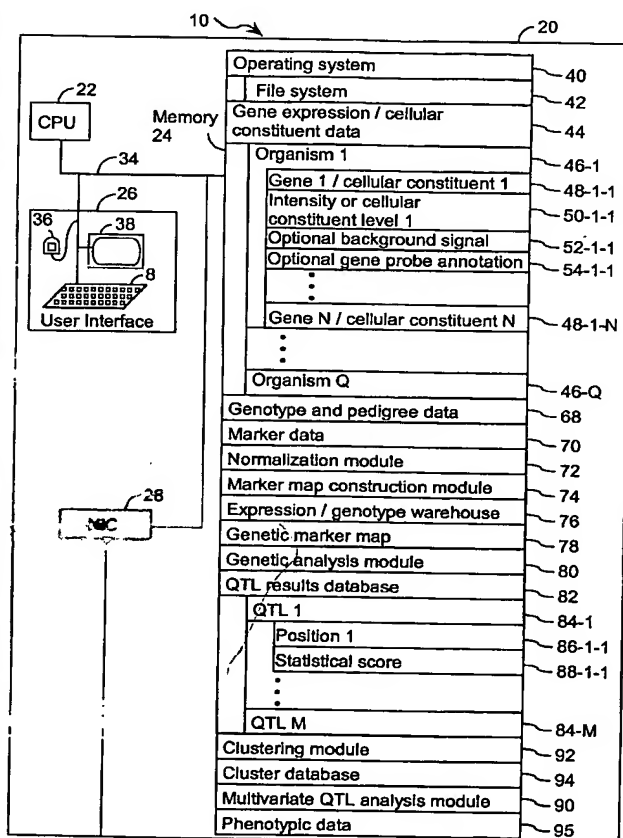
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- (71) Applicant (for all designated States except US):
ROSETTA INPHARMATICS LLC [US/US]; 12040
115th Avenue, N.E., Kirkland, WA 98034 (US).
- (72) Inventors; and
- (75) Inventors/Applicants (for US only): SCHADT, Eric [US/US]; 1517 3rd Place, Kirkland, WA 98033 (US).
MONKS, Stephanie, A. [US/US]; 906 N.E. 122nd Street, Seattle, WA 98125 (US).
- (74) Agents: ANTILER, Adriane, M. et al.; Pennie & Edmonds LLP, 1155 Avenue of the Americas, New York, NY 10036 (US).
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(54) Title: COMPUTER SYSTEMS AND METHODS FOR ASSOCIATING GENES WITH TRAITS USING CROSS SPECIES DATA



(57) Abstract: A method for confirming the association of a query QTL or a query gene in the genome of a second species with a clinical trait T exhibited by the second species. A first QTL or a first gene in a first species that is linked to a trait T' is found. The trait T' is indicative of trait T. A region of the genome of the first species that comprises the first QTL or the first gene is mapped to a particular region of the genome of the second species. A query QTL or a query gene in the second species that is potentially associated with the trait T is found. The potential association of the query QTL or the query gene with the clinical trait T is confirmed when the query QTL or the query gene is in the particular region of the genome of the second species.

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